

FIGURE 1

att ccg gct tct atg gag cac tcg gga cca ggt ccg cgg cgc gcg cac tcg ctc  
gct cgc cgc ccc cca gcc agc tct cgc ttc cgc gcc gcc agc cgc gcc ccg cgc  
ctc ctc gct gca ccc cgc gac cta gag cca aga aag ttt gtg tgg cga gtg agg  
gcc gga gag gag agc gcg ccc gcg gag tgc cgt cca gac cag cgc ggc ccc ggc  
gga gag ggg agc gcc ccg agc cca ggc ggc ggc ggc tag ccc gag tcc gcg acc  
ccc gcc cct ccg ccc gcc atg ggc gcc gcc gcc cgc agc ctg ccg ctc gcg ttc  
Cys Leu Leu Leu Leu Gly Thr Leu Leu Pro Arg Ala Asp Ala Cys Ser Cys Ser  
tgc ctc ctg ctg ctg ggg acg ctg ctc ccc cgg gcc gac gcc tgc agc tgc tcc  
Pro Val His Pro Gln Gln Ala Phe Cys Asn Ala Asp Ile Val Ile Arg Ala Lys  
ccg gtg cac ccg caa cag gcg ttt tgc aat gca gac ata gtg atc agg gcc aaa  
Ala Val Asn Lys Lys Glu Val Asp Ser Gly Asn Asp Ile Tyr Gly Asn Pro Ile  
gca gtc aat aag aag gag gtg gac tct ggc aac gac atc tac ggc aac ccc atc  
Lys Arg Ile Gln Tyr Glu Ile Lys Gln Ile Lys Met Phe Lys Gly Pro Asp Gln  
aag cgg att cag tat gag atc aag cag ata aag atg ttc aag gga cct gat cag  
Asp Ile Glu Phe Ile Tyr Thr Ala Pro Ala Ala Val Cys Gly Val Ser Leu  
gac ata gag ttt atc tac aca gcc ccc gcc gct gcc gtg tgt ggg gtc tcg ctg  
Asp Ile Gly Gly Lys Lys Glu Tyr Leu Ile Ala Gly Lys Ala Glu Gly Asn Gly  
gac att gga gga aag aag gag tat ctc att gca ggg aag gcc gag ggg aat ggc  
Asn Met His Ile Thr Leu Cys Asp Phe Ile Val Pro Trp Asp Thr Leu Ser Ala  
aat atg cat atc acc ctc tgt gac ttc atc gtg ccc tgg gac acc ctg agt gcc  
Thr Gln Lys Lys Ser Leu Asn His Arg Tyr Gln Met Gly Cys Glu Cys Lys Ile  
acc cag aag aag agc ctg aac cac agg tac cag atg ggc tgt gag tgc aag atc  
Thr Arg Cys Pro Met Ile Pro Cys Tyr Ile Ser Ser Pro Asp Glu Cys Leu Trp  
act cga tgc ccc atg atc cca tgc tac atc tcc tct ccg gac gag tgc ctc tgg  
Met Asp Trp Val Thr Glu Lys Asn Ile Asn Gly His Gln Ala Lys Phe Phe Ala  
atg gac tgg gtc acg gag aag aac atc aac gga cac cag gcc aag ttc ttc gcc  
Cys Ile Lys Arg Ser Asp Gly Ser Cys Ala Trp Tyr Arg Gly Ala Ala Pro Pro  
tgc atc aag aga agc gac ggc tcc tgc gcc tgg tac cgc gga gca gca ccc ccc  
Lys Gln Glu Phe Leu Asp Ile Glu Asp Pro  
aag cag gag ttt ctg gac atc gag gac ccg taa gca ggc cac cag gac tcc tgg  
ggc caa ttg aca gtg tcc aag agt tca gac tgg tcc agc tcc gac atc cct tcc  
tgg aca cag cat gaa taa a

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FIGURE 2

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att ccg gcc cgc cgt ccc oca ccc cgc cgc ccc gcc cgg cga att gcg ccc cgc  
 gcc cct ccc ctc gcg ccc ccg aga caa aga gga gag aaa gtt tgc gcg gcc gag  
 cgg ggc agg tga gga ggg tga gcc gcg cgg gag ggg ccc gcc tcg gcc ccg gct  
 cag ccc ccg ccc gcg ccc cca gcc cgc cgc cgc gag cag cgc ccg gac ccc cca  
 -26  
 gcg gcg gcc ccc gcc cgc cca gcc ccc cgg ccc gcc atg ggc gcc gcg gcc cgc  
 -20 Thr Leu Arg Leu Ala Leu Gly Leu Leu Leu -10 Ala Thr Leu Leu Arg Pro Ala  
 acc ctg cgg ctg gcg ctc ggc ctc ctg ctg ctg gcg acg ctg ctt cgc ccg gcc  
 -1 1 10  
 Asp Ala Cys Ser Cys Ser Pro Val His Pro Gln Gln Ala Phe Cys Asn Ala Asp  
 gac gcc tgc agc tgc tcc ccg gtg cac ccg caa cag gcg ttt tgc aat gca gat  
 20 30  
 Val Val Ile Arg Ala Lys Ala Val Ser Glu Lys Glu Val Asp Ser Gly Asn Asp  
 gta gtg atc agg gcc aaa gcg gtc agt gag aag gaa gtg gac tct gga aac gac  
 40 50  
 Ile Tyr Gly Asn Pro Ile Lys Arg Ile Gln Tyr Glu Ile Lys Gln Ile Lys Met  
 att tat ggc aac cct atc aag agg atc cag tat gag atc aag cag ata aag atg  
 60 70  
 Phe Lys Gly Pro Glu Lys Asp Ile Glu Phe Ile Tyr Thr Ala Pro Ser Ser Ala  
 ttc aaa ggg cct gag aag gat ata gag ttt atc tac acg gcc ccc tcc tcg gca  
 80 90  
 Val Cys Gly Val Ser Leu Asp Val Gly Gly Lys Lys Glu Tyr Leu Ile Ala Gly  
 gtg tgt ggg gtc tcg ctg gac gtt gga gga aag aag gaa tat ctc att gca gga  
 100  
 Lys Ala Glu Gly Asp Gly Lys Met His Ile Thr Leu Cys Asp Phe Ile Val Pro  
 aag gcc gag ggg gac ggc aag atg cac atc acc ctc tgt gac ttc atc gtg ccc  
 110 120  
 Trp Asp Thr Leu Ser Thr Thr Gln Lys Lys Ser Leu Asn His Arg Tyr Gln Met  
 tgg gac acc ctg agc acc acc cag aag aag agc ctg aac cac agg tac cag atg  
 130 140  
 Gly Cys Glu Cys Lys Ile Thr Arg Cys Pro Met Ile Pro Cys Tyr Ile Ser Ser  
 ggc tgc gag tgc aag atc acg cgc tgc ccc atg atc ccg tgc tac atc tcc tcc  
 150 160  
 Pro Asp Glu Cys Leu Trp Met Asp Trp Val Thr Glu Lys Asn Ile Asn Gly His  
 ccg gac gag tgc ctc tgg atg gac tgg gtc aca gag aag aac atc aac ggg cac  
 170 180  
 Gln Ala Lys Phe Phe Ala Cys Ile Lys Arg Ser Asp Gly Ser Cys Ala Trp Tyr  
 cag gcc aag ttc ttc gcc tgc atc aag aga agt gac ggc tcc tgt gcg tgg tac  
 190 194  
 Arg Gly Ala Ala Pro Pro Lys Gln Glu Phe Leu Asp Ile Glu Asp Pro  
 cgc ggc gcg gcg ccc ccc aag cag gag ttt ctc gac atc gag gac cca taa gca  
 ggc ctc caa cgc ccc tgt ggc caa ctg caa aaa aag cct cca agg gtt tcg act  
 ggt cca gct ctg aca tcc ctt cct gga aac agc atg aat aaa aca ctc atc ccc  
 gga att c

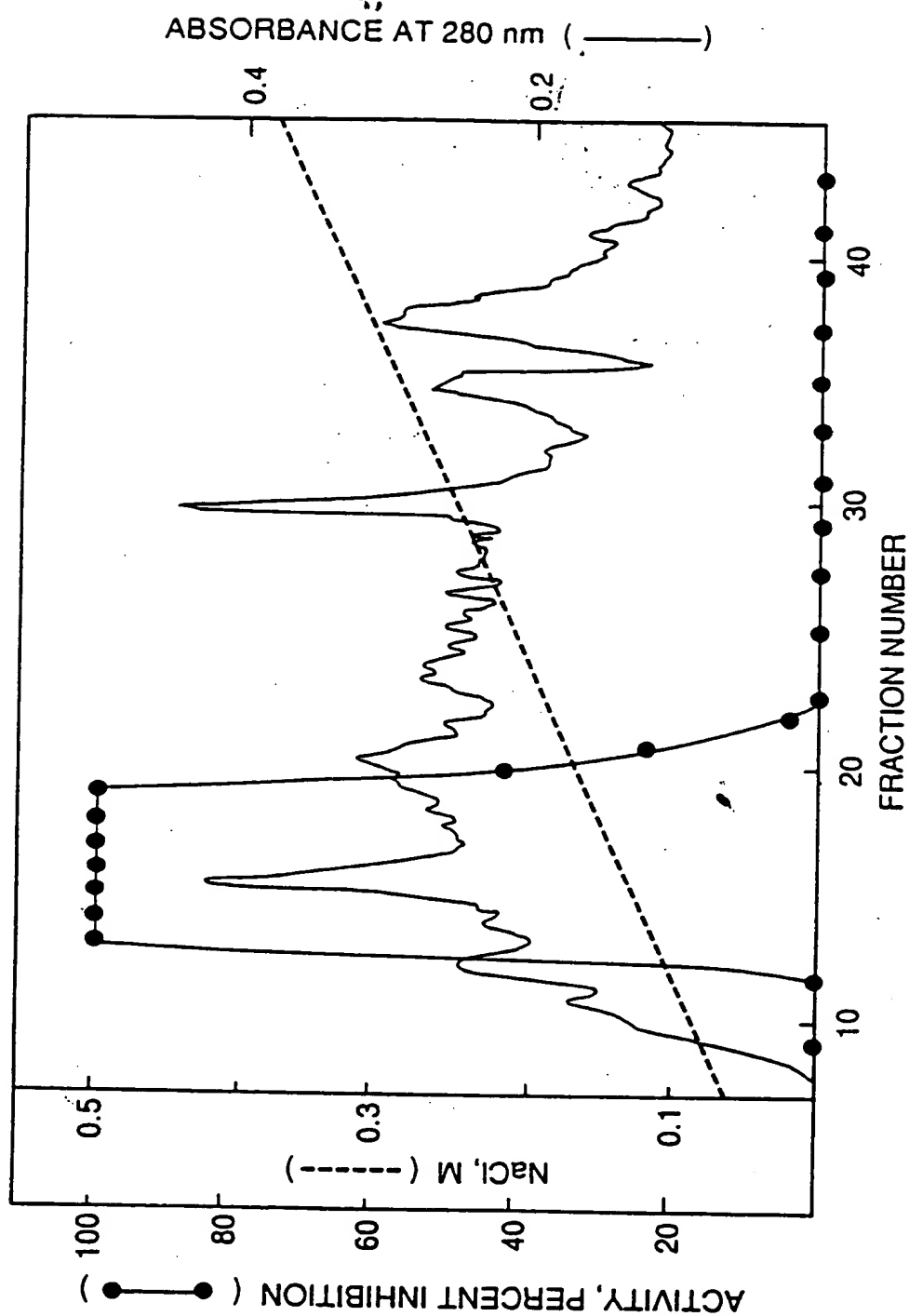
261220 456E0880

FIGURE 3

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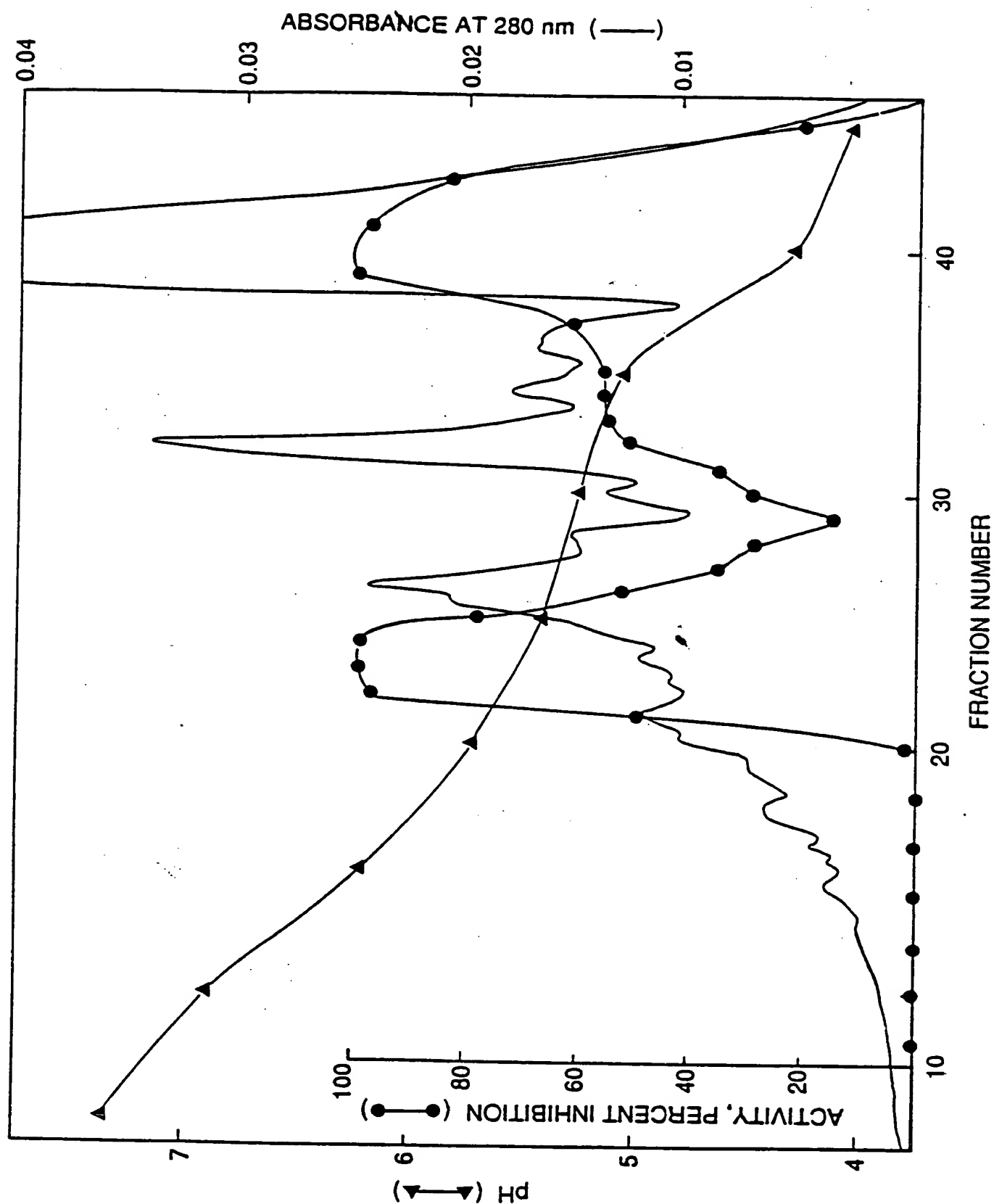
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FIGURE 4

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FIGURE 5

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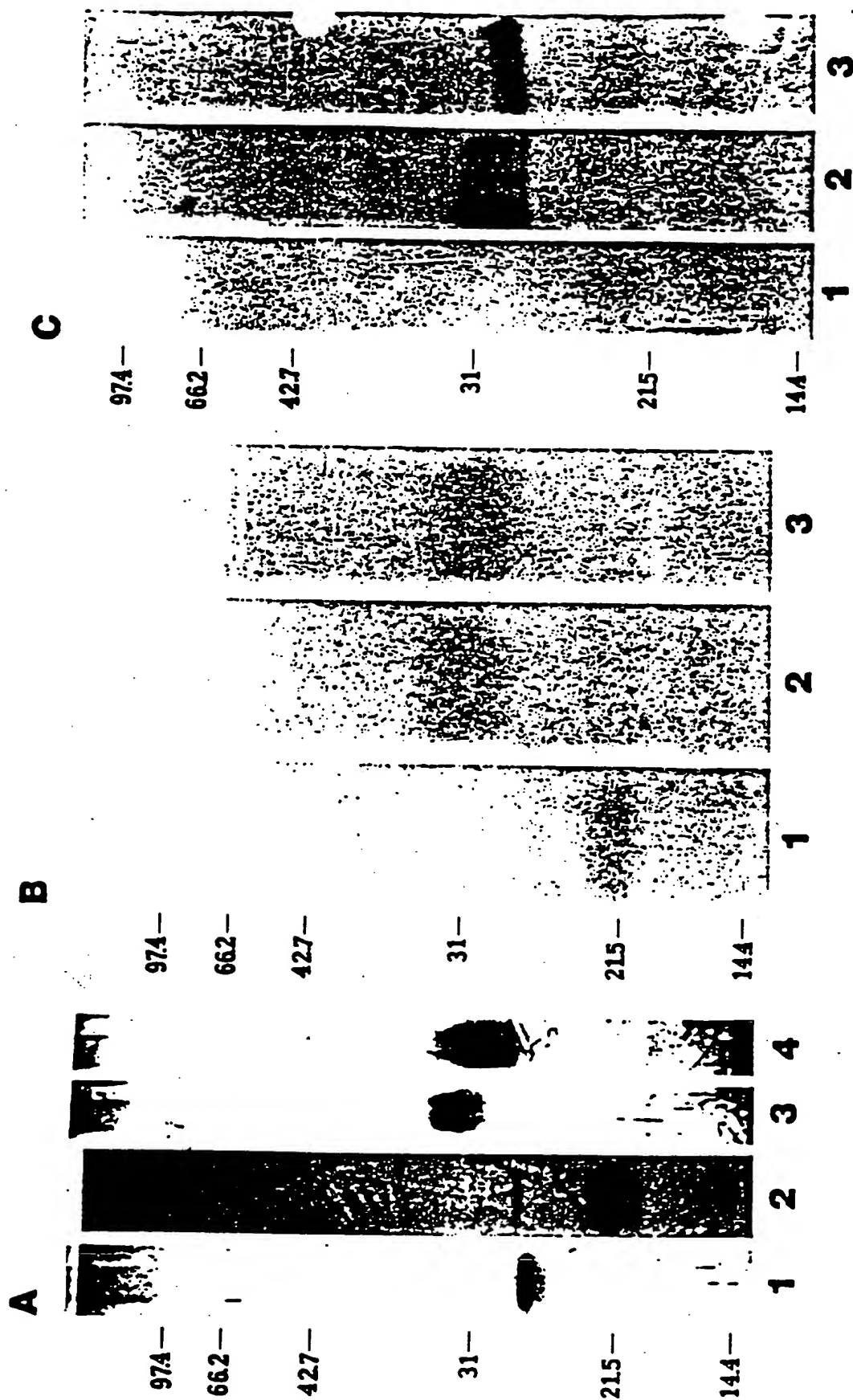


FIGURE 6

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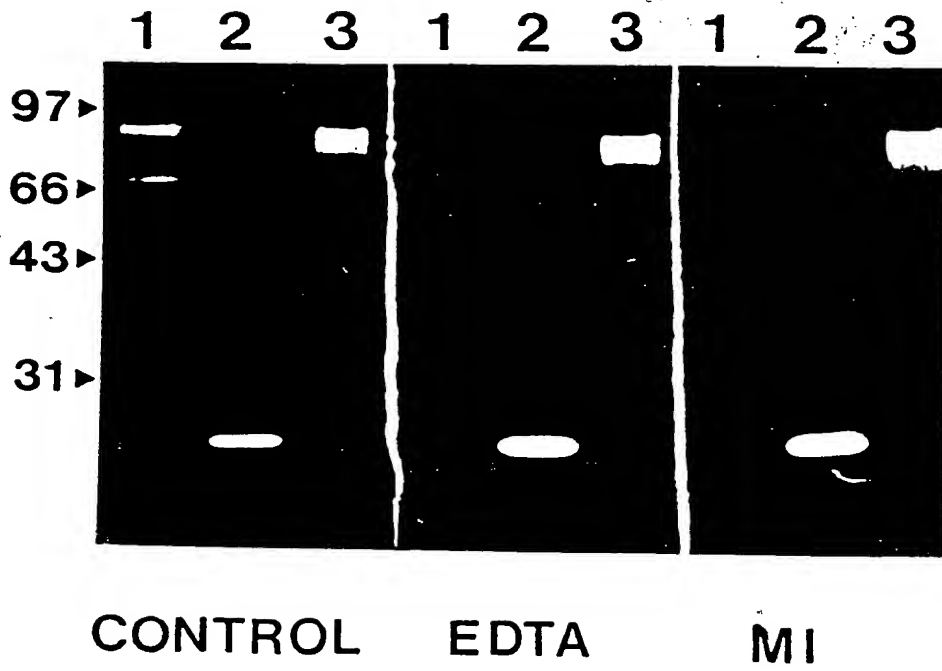
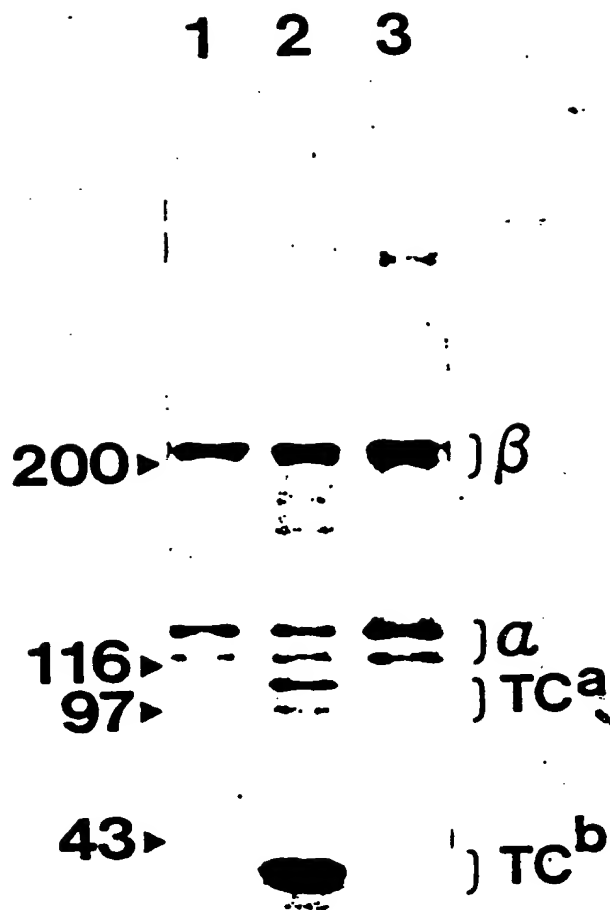


FIGURE 7



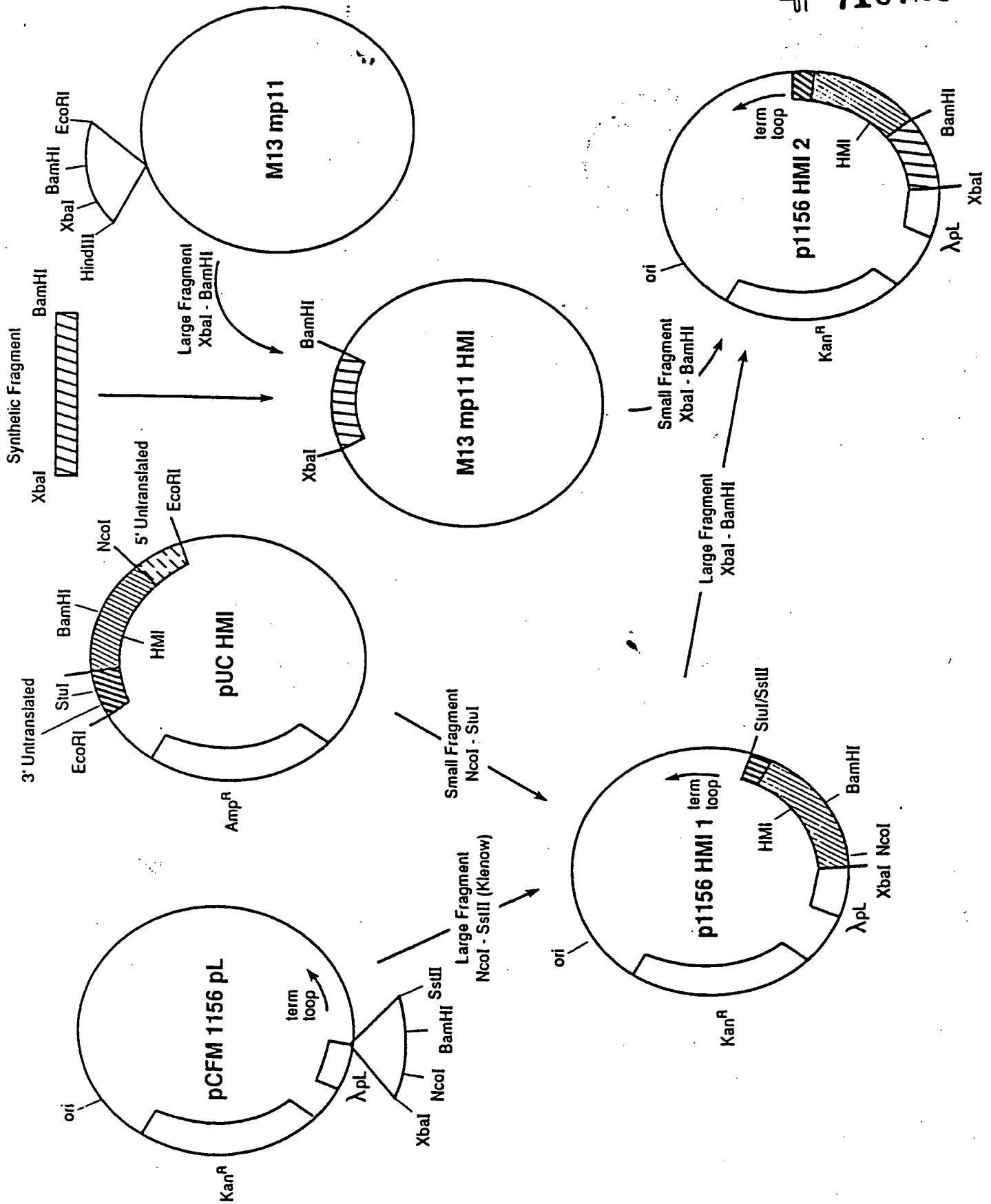
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FIGURE 8

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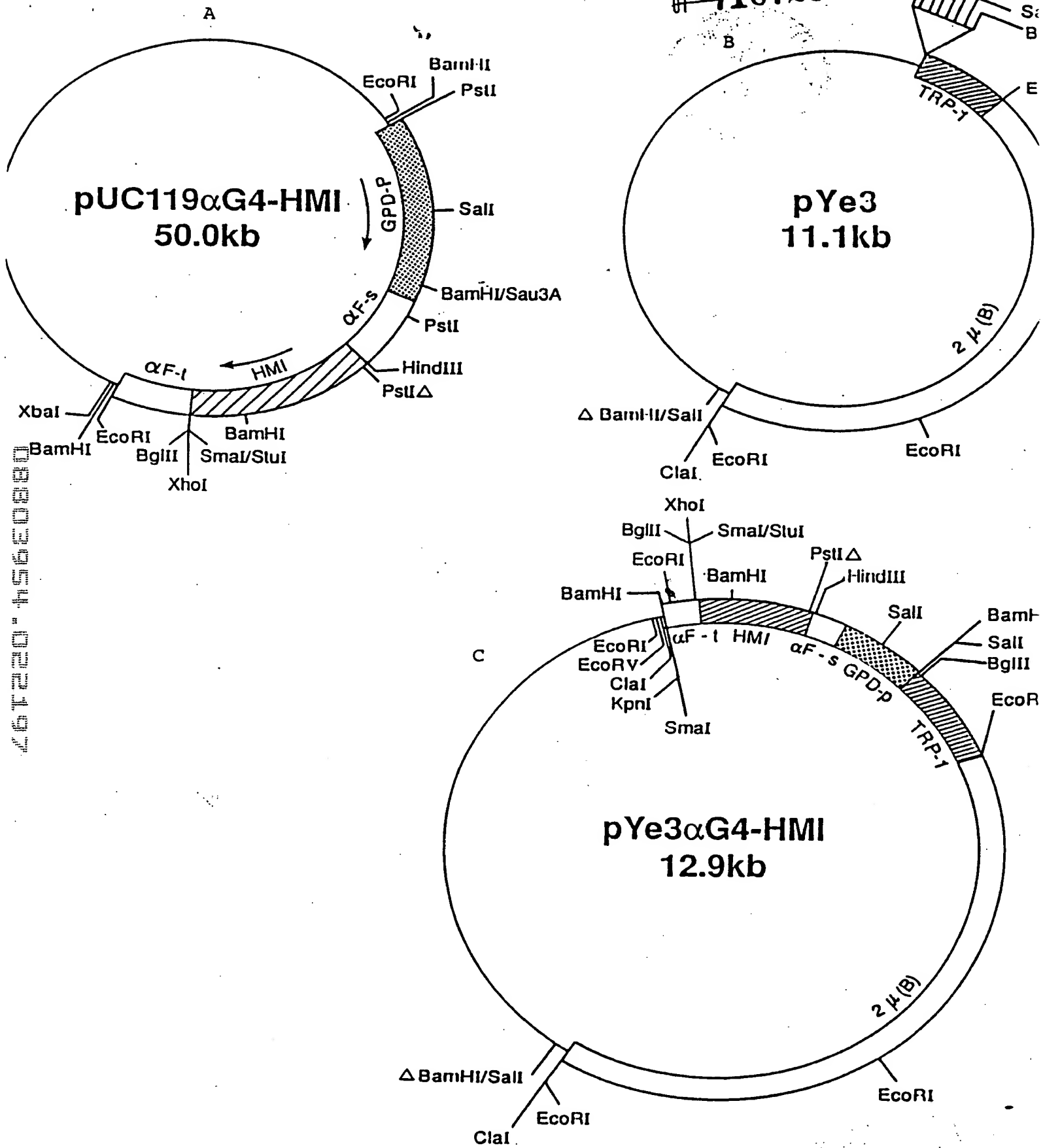
FIGURE 9



|            |            |            |             |            |            |
|------------|------------|------------|-------------|------------|------------|
| 10         | 20         | 30         | 40          | 50         | 60         |
| CTAGAAAAAA | CCAAGGAGGT | AATAAATAAT | GTGTTCTTGT  | TCTCCTGTAC | ACCCTCAACA |
| TTTTTT     | GGTTCCTCCA | TTATTTATTA | CACAAGAACA  | AGAGGACATG | TGGGAGTTGT |
| 70         | 80         | 90         | 100         | 110        | 120        |
| AGCTTTTTGT | AACGCTGATG | TAGTTATCCG | TGCAAAAGCT  | GTTTCTGAAA | AAGAAGTTGA |
| TCGAAAAACA | TTGCGACTAC | ATCAATAGGC | ACGTTTTTCGA | CAAAGACTTT | TTCTTCAACT |
| 130        | 140        | 150        | 160         |            |            |
| TTCTGGTAAC | GACATCTACG | GTAACCCGAT | CAAAAG.     |            |            |
| AAGACCATTG | CTGTAGATGC | CATTGGGCTA | GTTTTCCTAG  |            |            |

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FIGURE 10

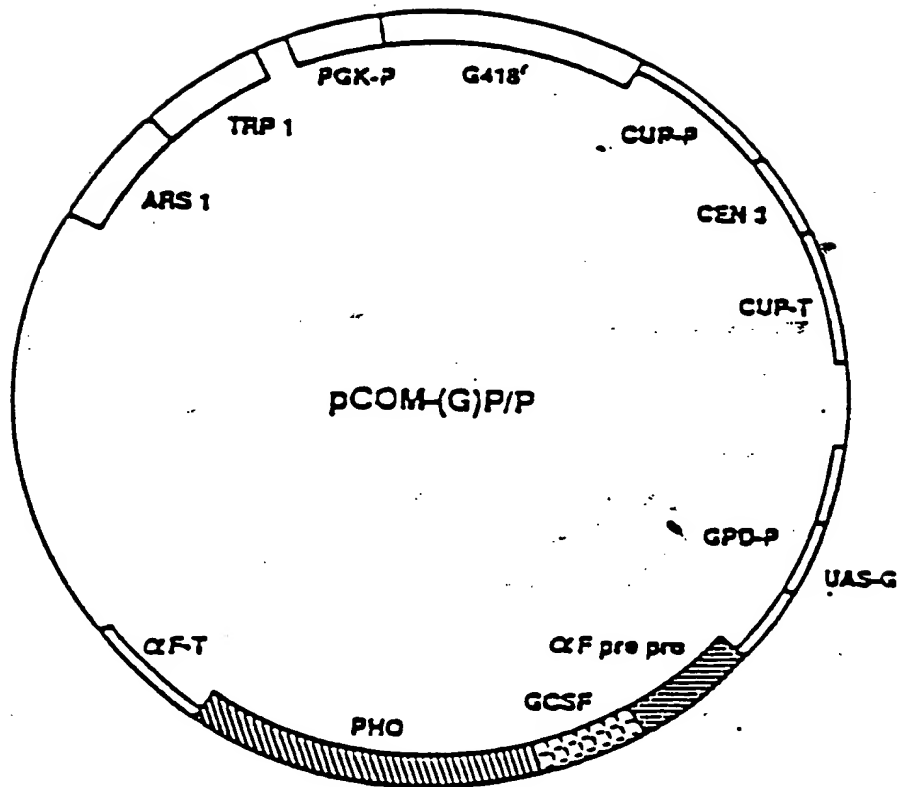


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FIGURE 11

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FIGURE 12

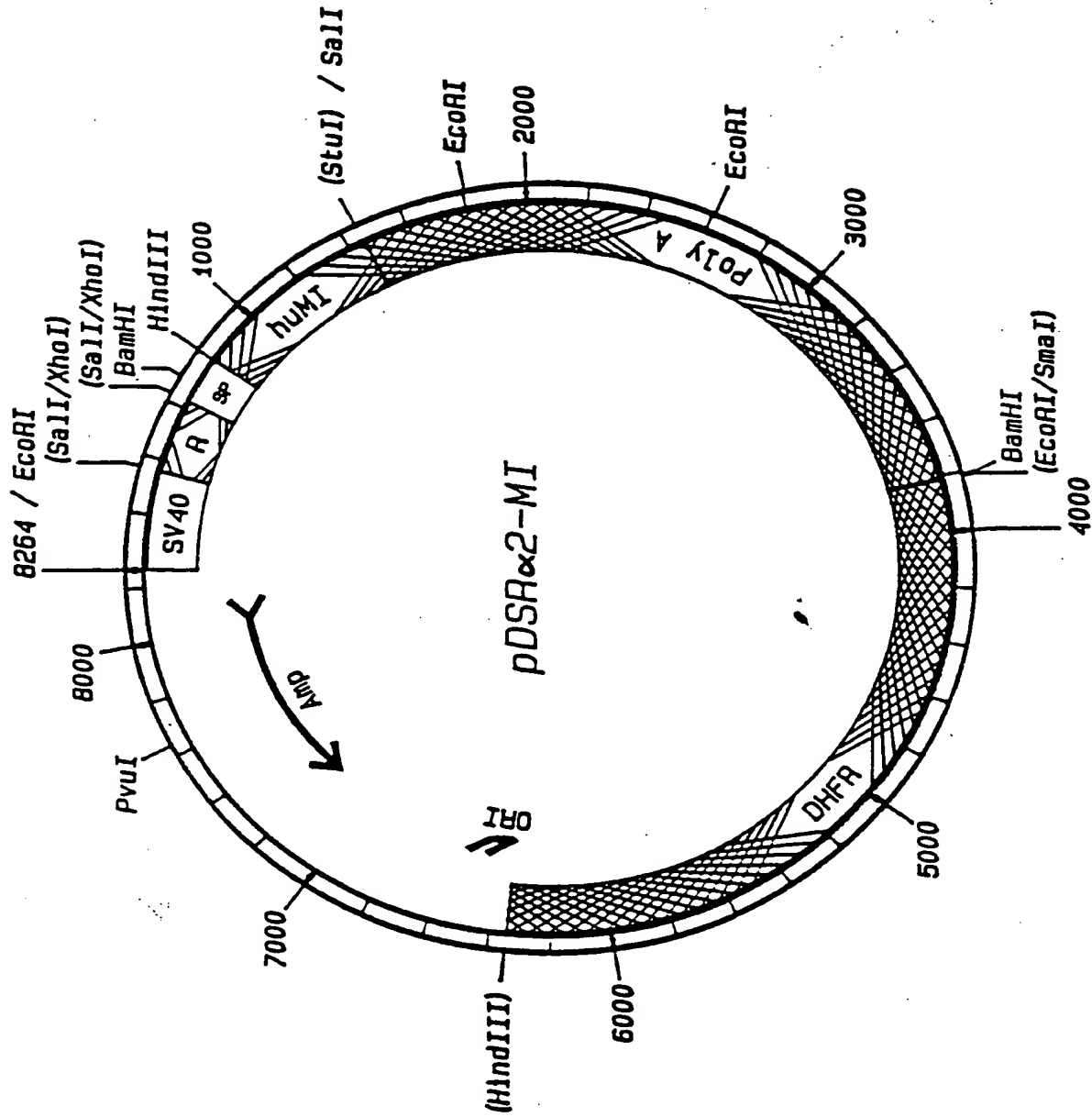


FIGURE 13

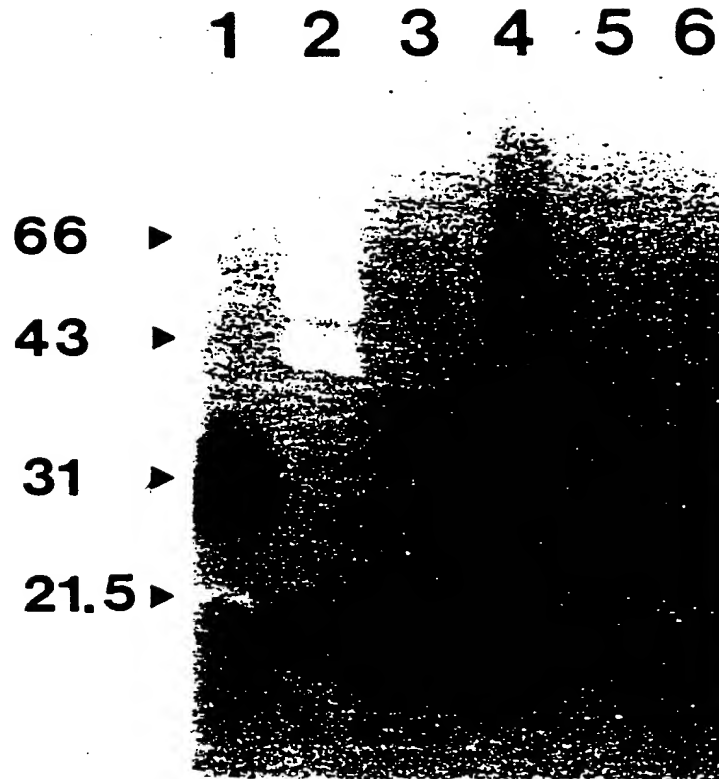
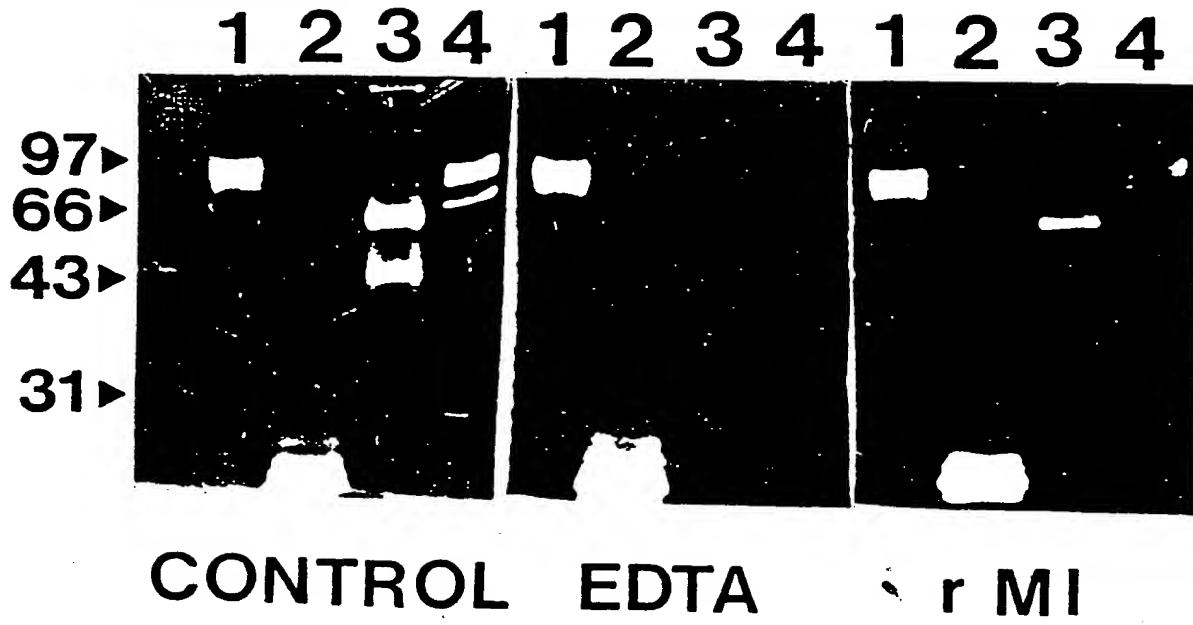


FIGURE 14



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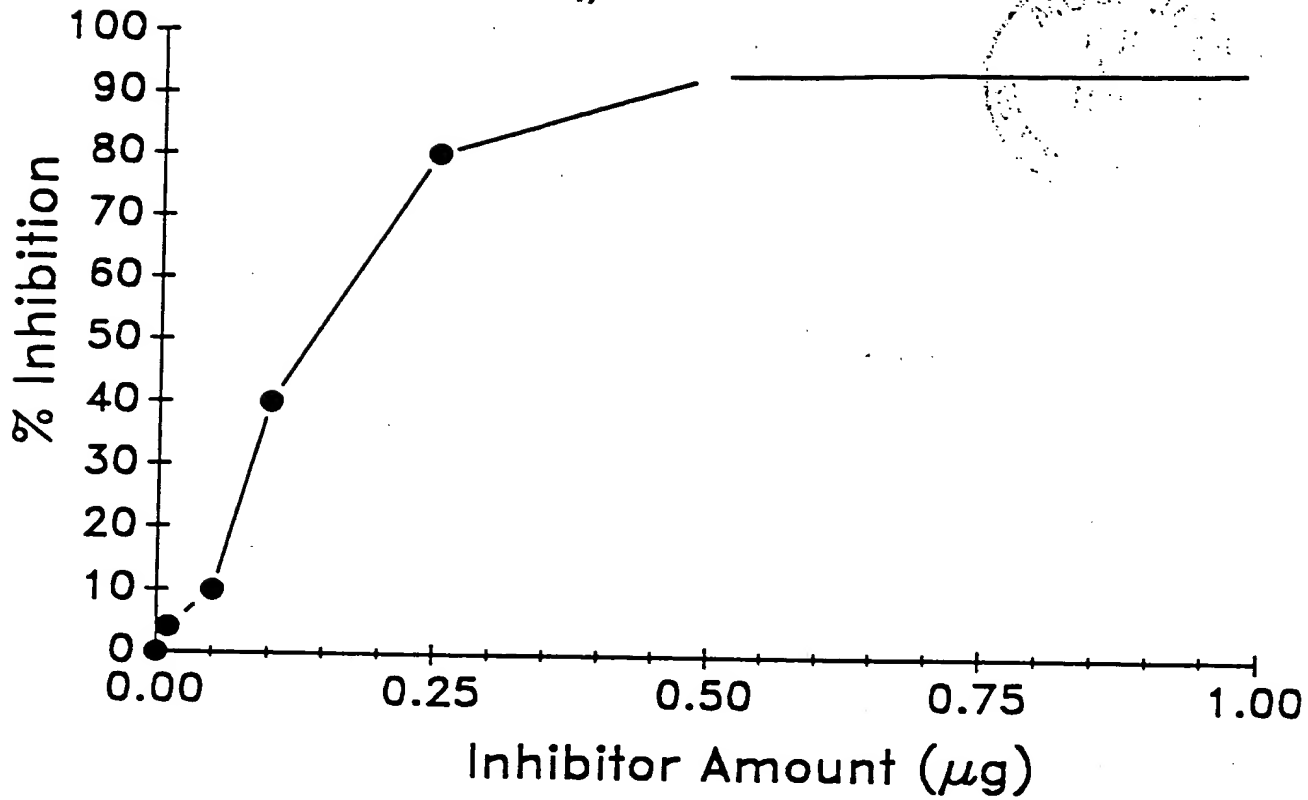
FIGURE 15

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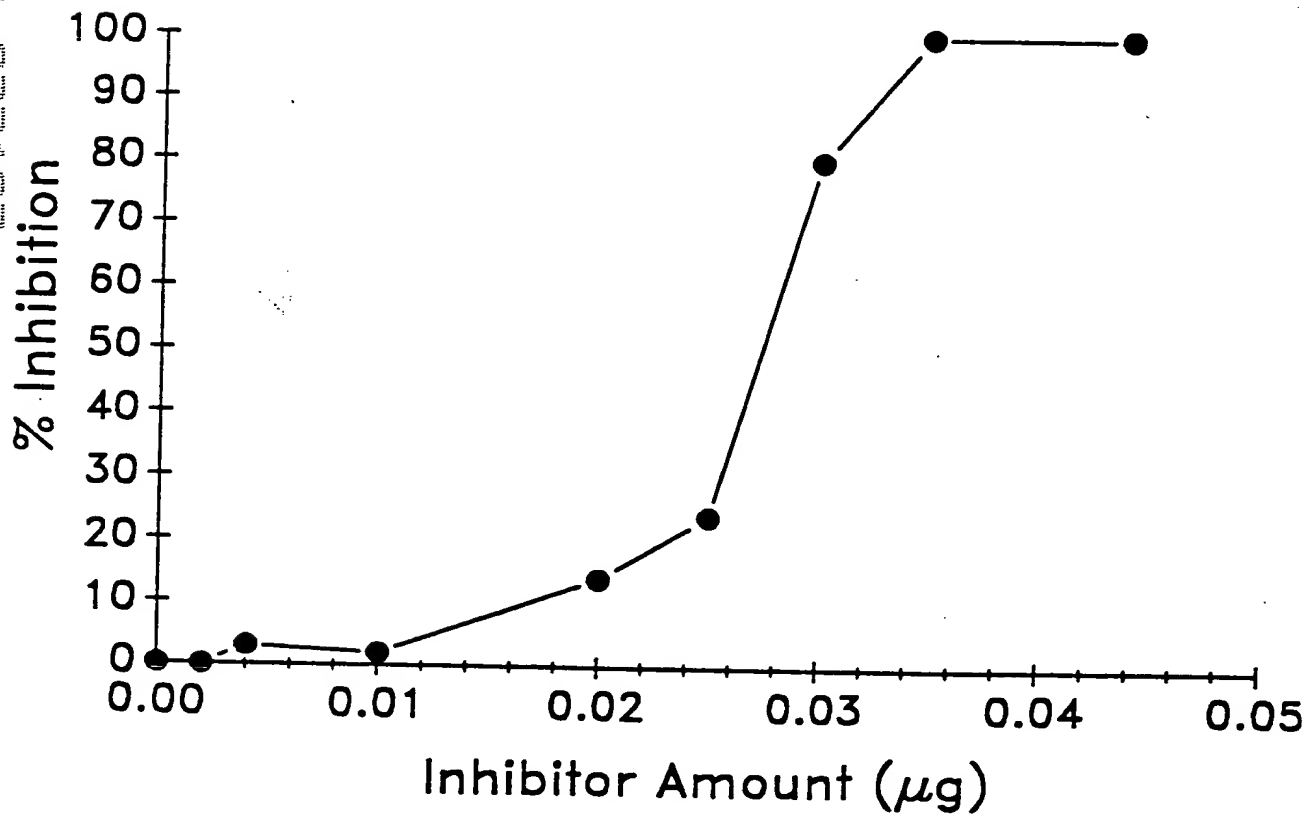
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A.



B.



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FIGURE 16

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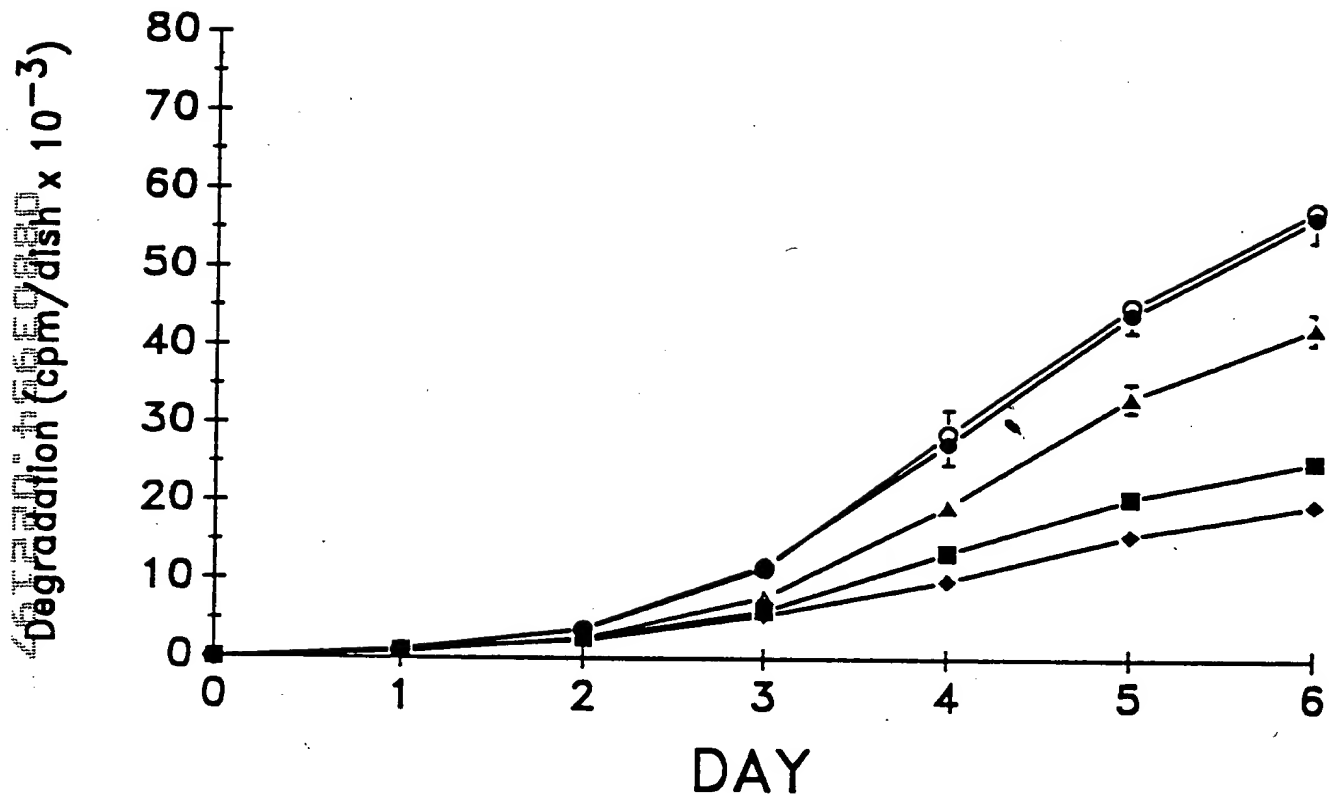




FIGURE 17

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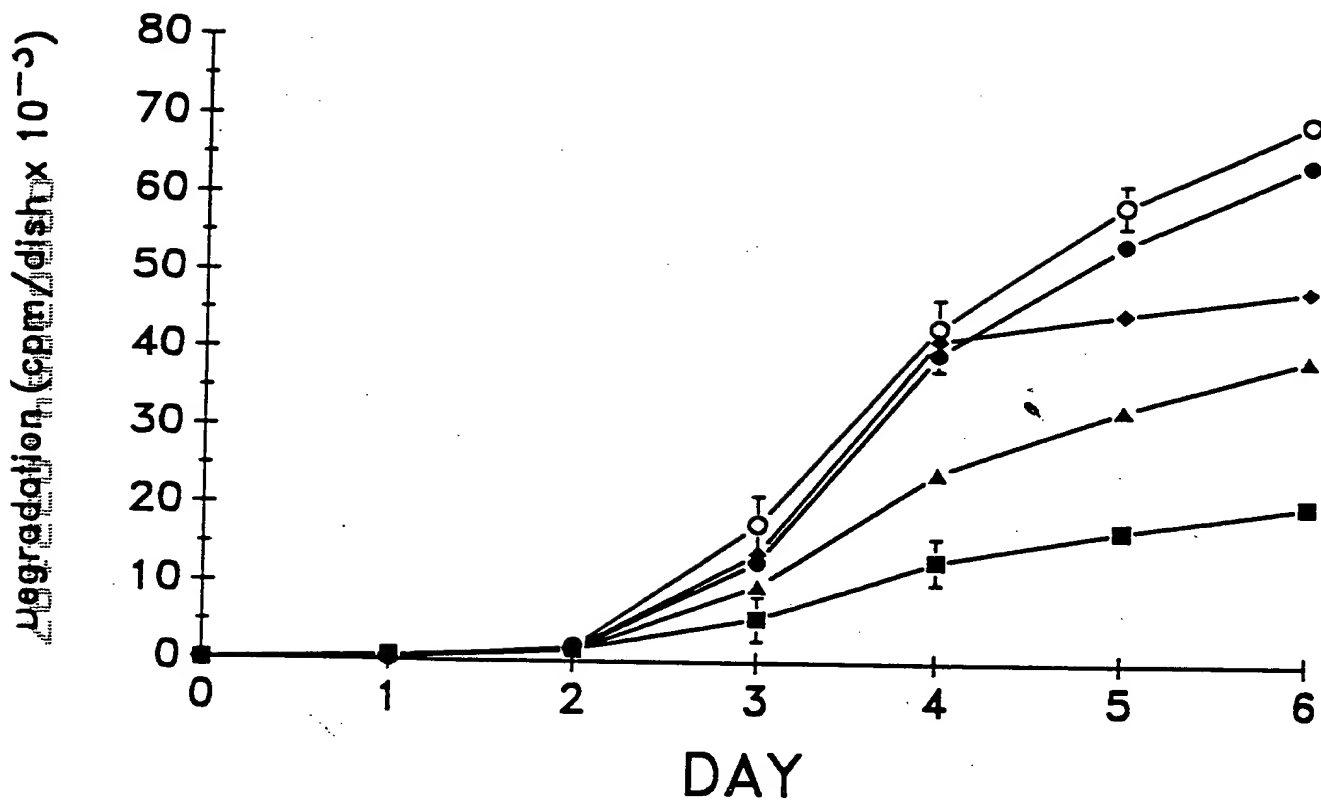
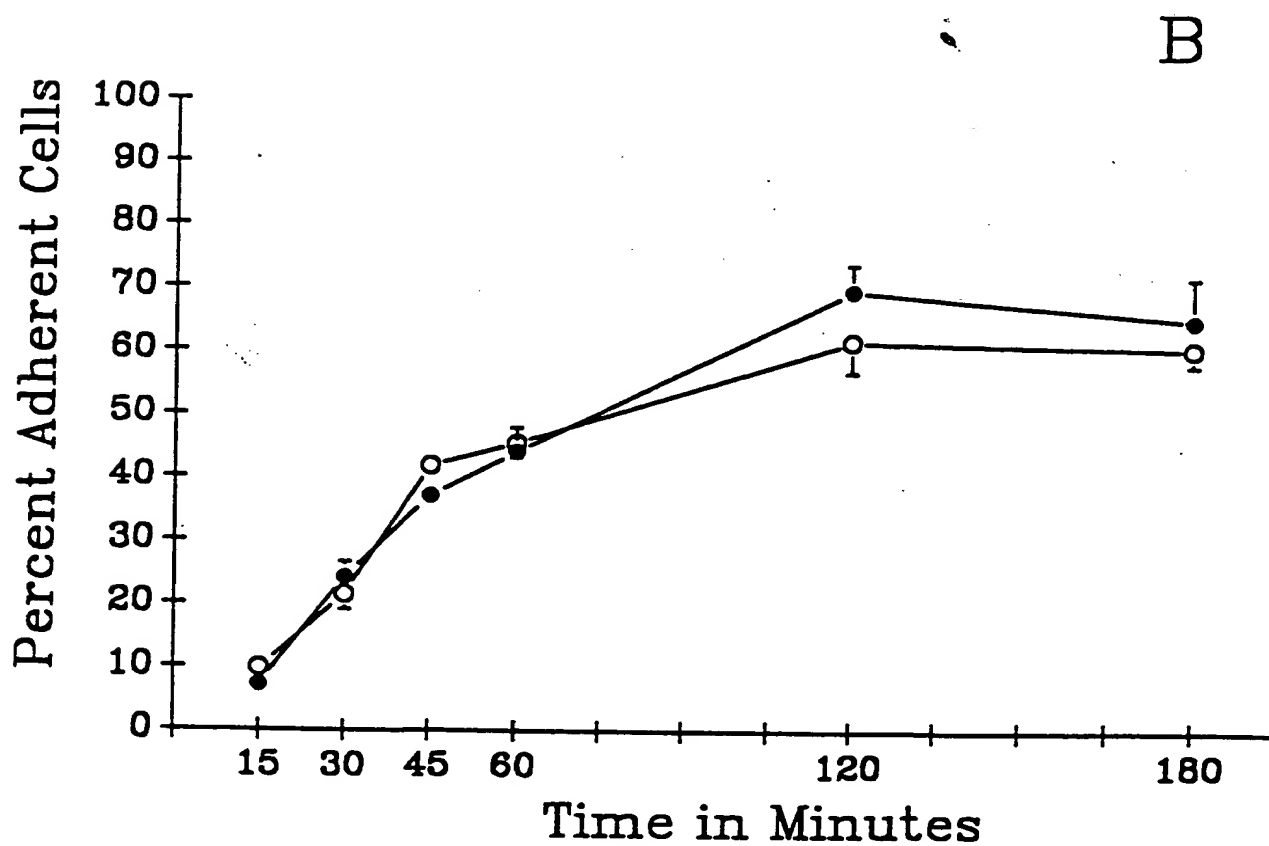
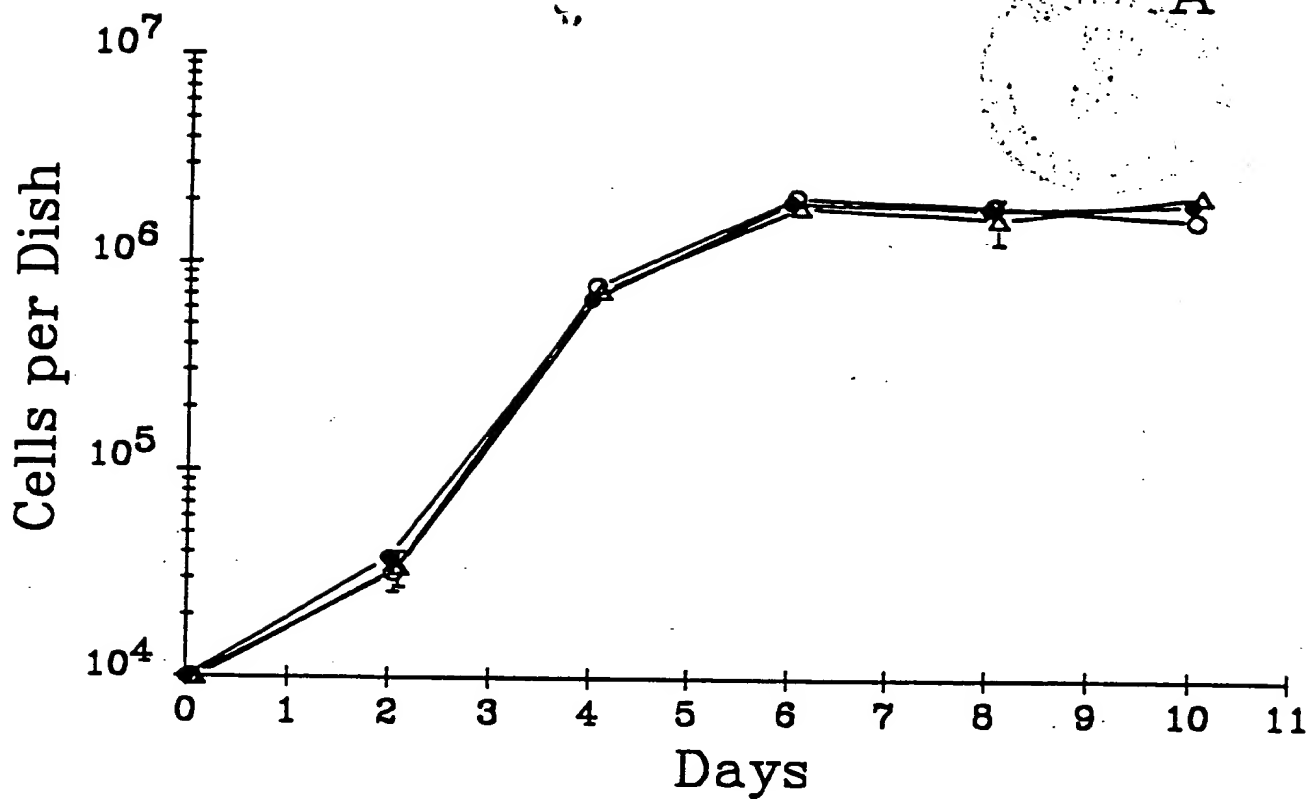


FIGURE 18

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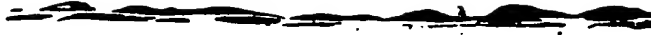
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FIGURE 19

A



B



C



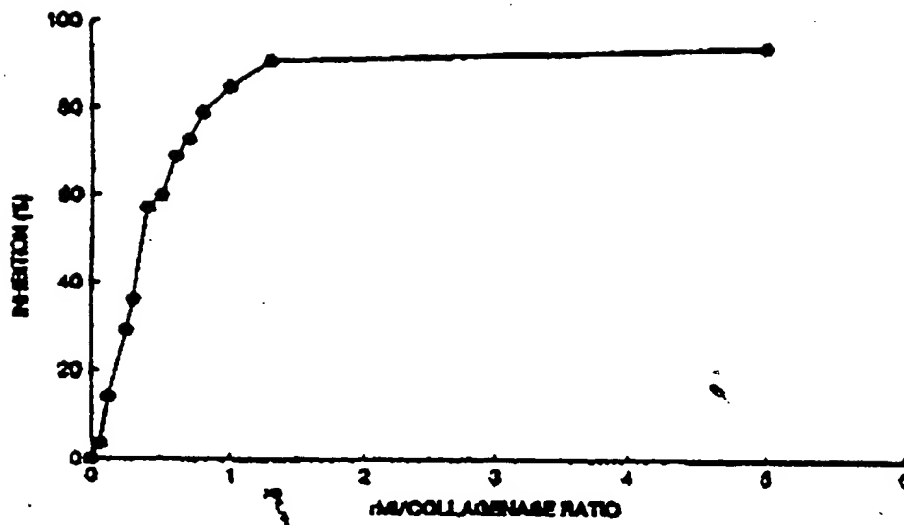
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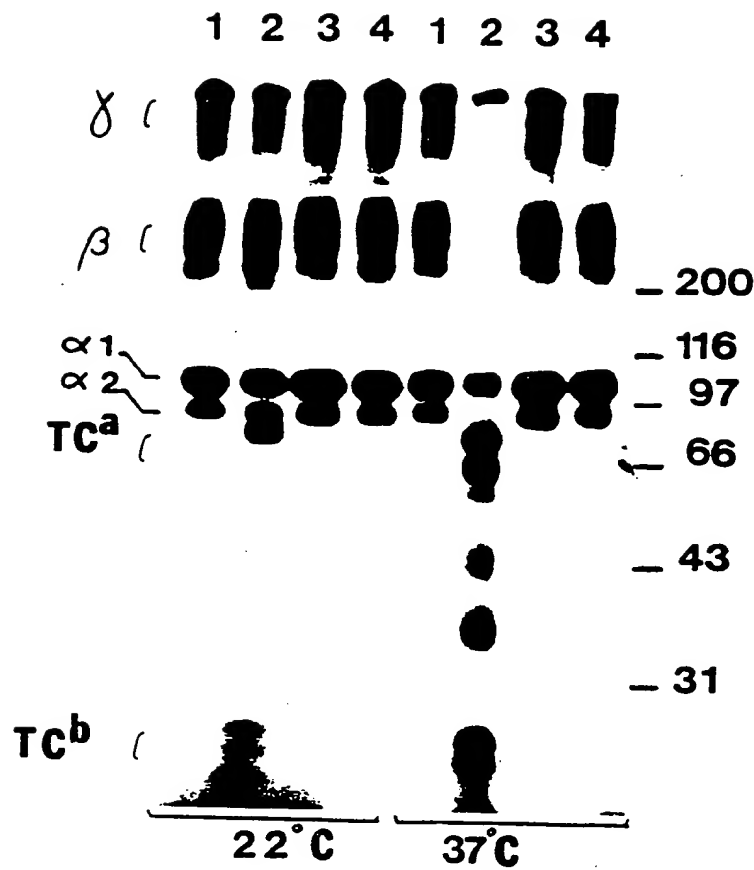
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FIGURE 20



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FIGURE 21



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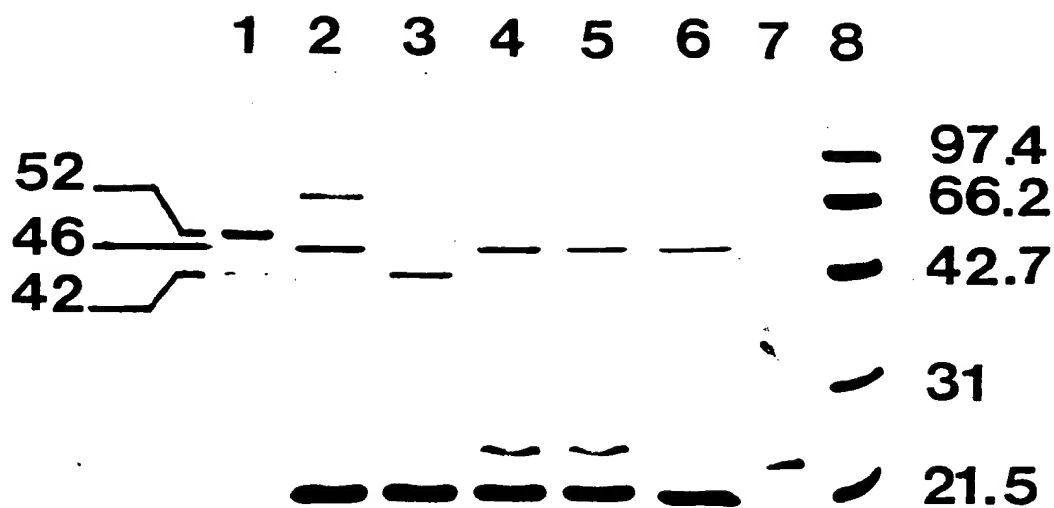
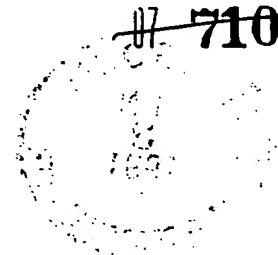
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FIGURE 23

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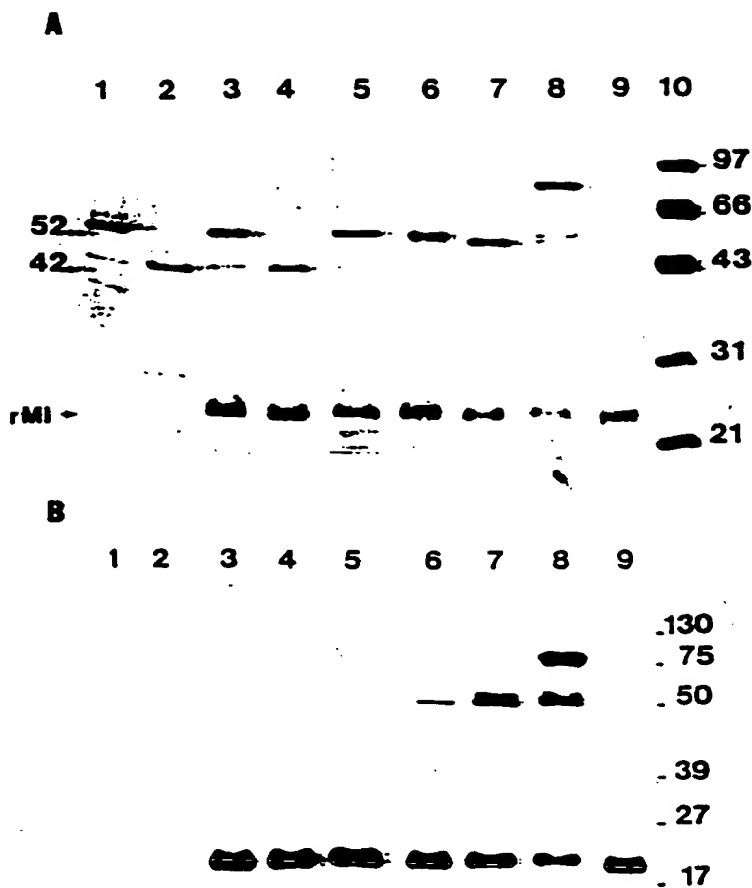
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FIGURE 24



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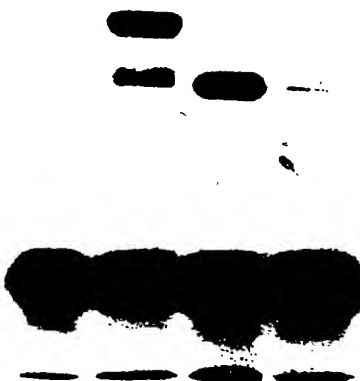
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FIGURE 25

1 2 3 4

procoll-rMI ▶  
coll-rMI ▶

rMI ▶

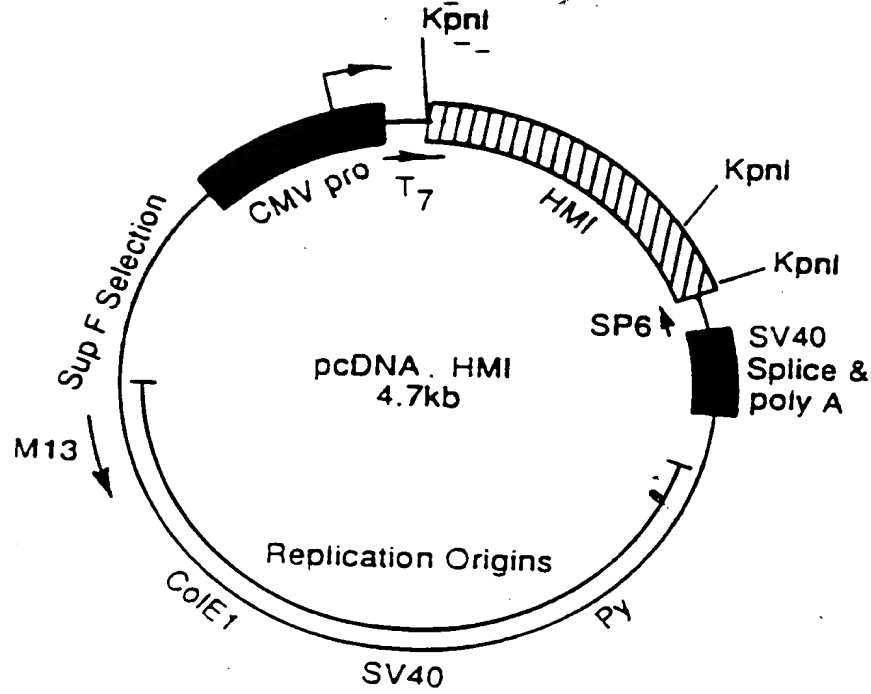


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FIGURE 26

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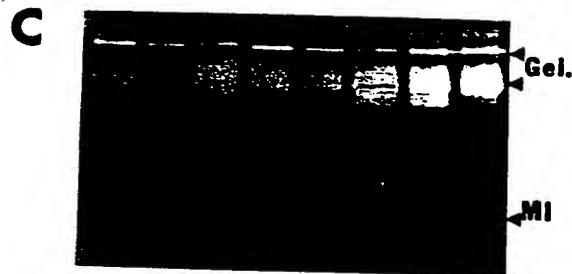
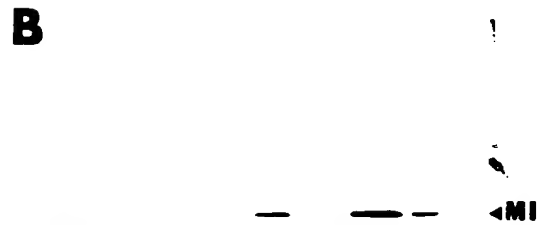
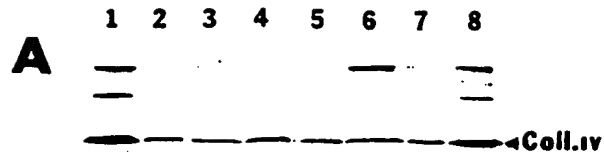
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FIGURE 27



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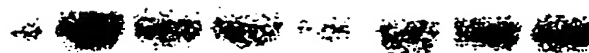
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FIGURE 28

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1 2 3 4 5 6 7 8 6 7



-28s



-18s

Total

Poly A

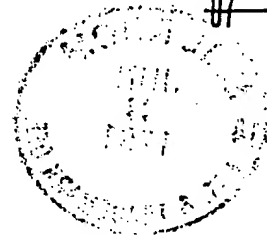
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FIGURE 29

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4R 8.60m1 m2 m3 m4 m5

A

B

n1 n2 n3 n4 n5

C

185

13

M1 8.60 m1 m2 m3 m4 m5

D

75

50

39

27

E

Gel

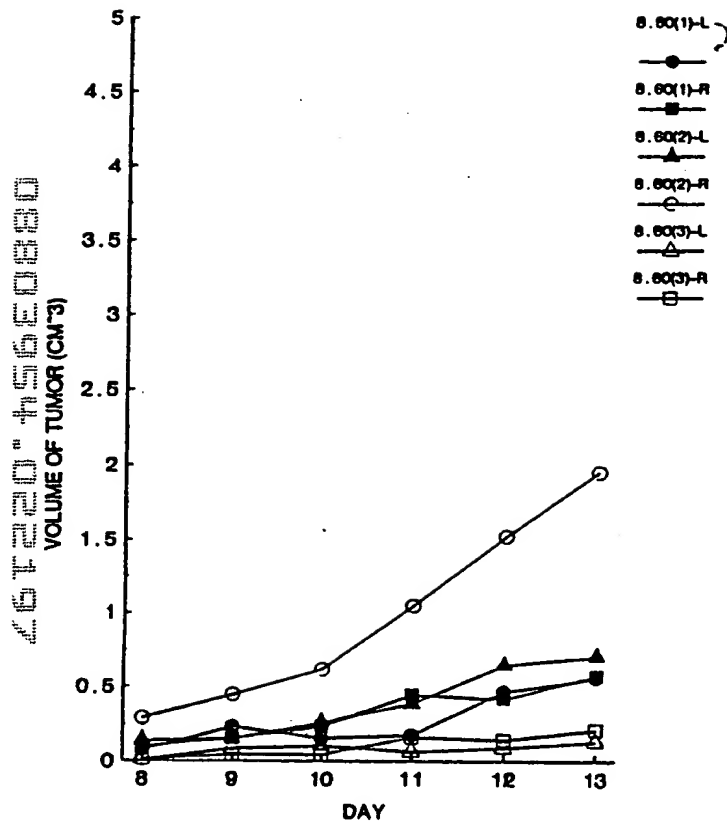
M1

0803954 0219

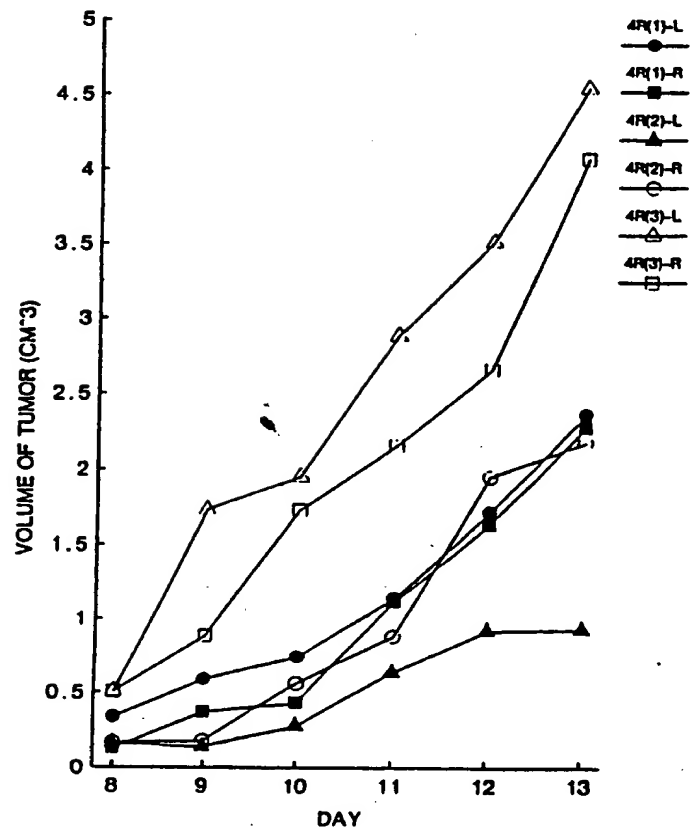
FIGURE 30

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A



B

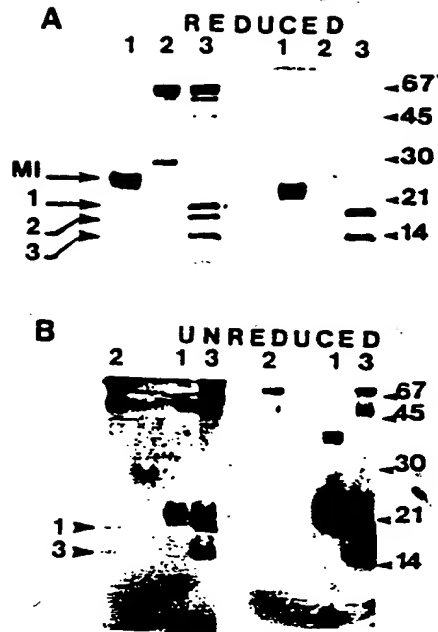


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FIGURE 31



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FIGURE 32

Band 1: (Cys)-Ser-(Cys)-Ser-Pro-Val-His-Pro-Gln-Gln-Ala-Phe-  
(Cys)-Asn-Ala-Asp-Val-Val-Ile-....  
Band 2: Val-Val-Gly-Gly-Pro-(or Ala)-Val-Ala-His-Pro-His-Ser-  
Trp-Pro-Thr-Gln-Val-Ser-Leu-Arg-Thr-....  
Band 3: (Cys)-Ser-(Cys)-Ser-Pro-Val-....

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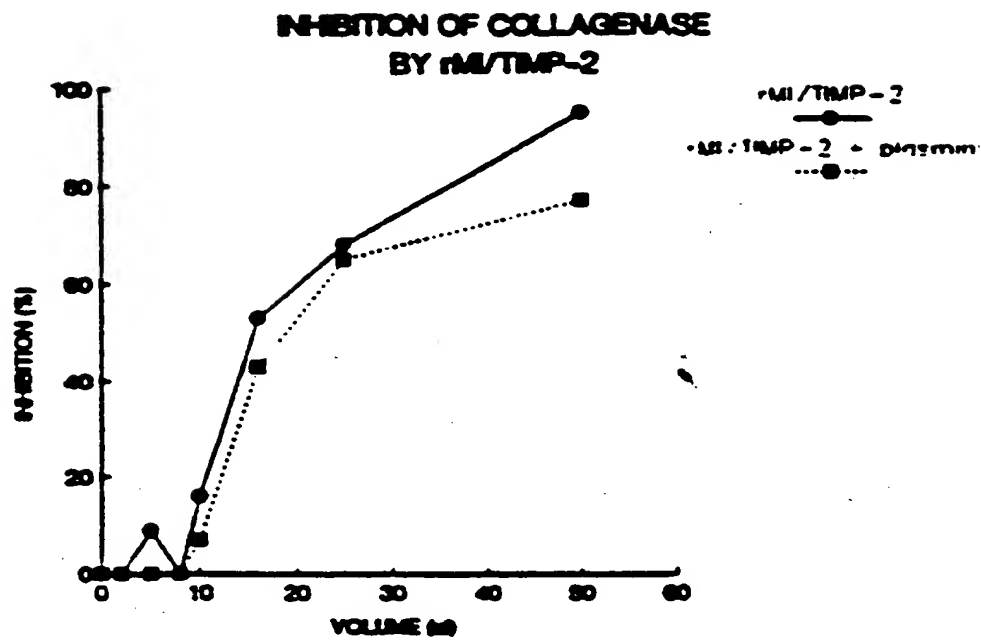


FIGURE 33

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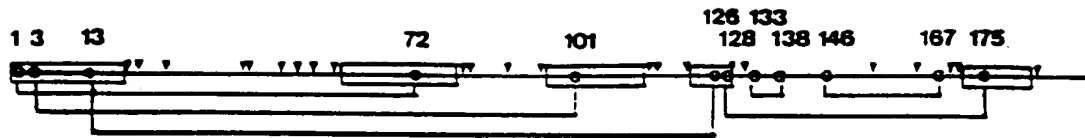
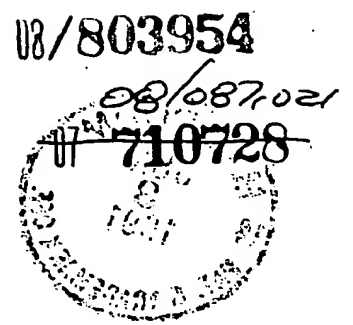
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FIGURE 34



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